

ABSTRACT OF THE DISCLOSURE

A method of identifying and archiving a nucleic acid sequence includes a) creating a directory of files in a computer, for storing information related to the nucleic acid sequence; b) inputting a raw nucleic acid sequence into the computer; c) trimming the raw nucleic acid sequence to obtain a trimmed nucleic acid sequence; d) submitting the trimmed nucleic acid sequence electronically to a nucleic acid identification database having a search program and receiving search results electronically from the nucleic acid identification database; e) choosing selective information from each search result and inserting the selective information from each search result into a first electronic spreadsheet; and f) selecting at least one of the search results from the first electronic spreadsheet and inserting the at least one search result into a second electronic spreadsheet.